

## RAW SEQUENCE LISTING

met/

**The Biotechnology Systems Branch of the Scientific and Technical  
Information Center (STIC) no errors detected.**

Application Serial Number: 10/565,495  
Source: IFWP  
Date Processed by STIC: 1/30/06

# ***ENTERED***

**RAW SEQUENCE LISTING** DATE: 01/30/2006  
PATENT APPLICATION: US/10/565,495 TIME: 15:09:22

Input Set : A:\PU60406SEQLIST.txt  
Output Set : N:\CRP4\01302006\J565495.raw

file://C:\CRF4\OUTHOLD\VsJ565495.htm

1/30/2006

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DATE: 01/30/2006

PATENT APPLICATION: US/10/565,495

TIME: 15:09:22

Input Set : A:\PU60406SEQLIST.txt

Output Set: N:\CRF4\01302006\J565495.raw

58 &lt;400&gt; SEQUENCE: 2

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59 Met Ser Leu Trp Leu Gly Ala Pro Val Pro Asp Ile Pro Pro Asp Ser
60 1 5 10 15
61 Ala Val Glu Leu Trp Lys Pro Gly Ala Gln Asp Ala Ser Ser Gln Ala
62 20 25 30
63 Gln Gly Gly Ser Ser Cys Ile Leu Arg Glu Glu Ala Arg Met Pro His
64 35 40 45
65 Ser Ala Gly Gly Thr Ala Gly Val Gly Leu Glu Ala Ala Glu Pro Thr
66 50 55 60
67 Ala Leu Leu Thr Arg Ala Glu Pro Pro Ser Glu Pro Thr Glu Ile Arg
68 65 70 75 80
69 Pro Gln Lys Arg Lys Lys Gly Pro Ala Pro Lys Met Leu Gly Asn Glu
70 85 90 95
71 Leu Cys Ser Val Cys Gly Asp Lys Ala Ser Gly Phe His Tyr Asn Val
72 100 105 110
73 Leu Ser Cys Glu Gly Cys Lys Gly Phe Phe Arg Arg Ser Val Ile Lys
74 115 120 125
75 Gly Ala His Tyr Ile Cys His Ser Gly Gly His Cys Pro Met Asp Thr
76 130 135 140
77 Tyr Met Arg Arg Lys Cys Gln Glu Cys Arg Leu Arg Lys Cys Arg Gln
78 145 150 155 160
79 Ala Gly Met Arg Glu Glu Cys Val Leu Ser Glu Glu Gln Ile Arg Leu
80 165 170 175
81 Lys Lys Leu Lys Arg Gln Glu Glu Glu Gln Ala His Ala Thr Ser Leu
82 180 185 190
83 Pro Pro Arg Arg Ser Ser Pro Pro Gln Ile Leu Pro Gln Leu Ser Pro
84 195 200 205
85 Glu Gln Leu Gly Met Ile Glu Lys Leu Val Ala Ala Gln Gln Gln Cys
86 210 215 220
87 Asn Arg Arg Ser Phe Ser Asp Arg Leu Arg Val Thr Pro Trp Pro Met
88 225 230 235 240
89 Ala Pro Asp Pro His Ser Arg Glu Ala Arg Gln Gln Arg Phe Ala His
90 245 250 255
91 Phe Thr Glu Leu Ala Ile Val Ser Val Gln Glu Ile Val Asp Phe Ala
92 260 265 270
93 Lys Gln Leu Pro Gly Phe Leu Gln Leu Ser Arg Glu Asp Gln Ile Ala
94 275 280 285
95 Leu Leu Lys Thr Ser Ala Ile Glu Val Met Leu Leu Glu Thr Ser Arg
96 290 295 300
97 Arg Tyr Asn Pro Gly Ser Glu Ser Ile Thr Phe Leu Lys Asp Phe Ser
98 305 310 315 320
99 Tyr Asn Arg Glu Asp Phe Ala Lys Ala Gly Leu Gln Val Glu Phe Ile
100 325 330 335
101 Asn Pro Ile Phe Glu Phe Ser Arg Ala Met Asn Glu Leu Gln Leu Asn
102 340 345 350
103 Asp Ala Glu Phe Ala Leu Leu Ile Ala Ile Ser Ile Phe Ser Ala Asp
104 355 360 365
105 Arg Pro Asn Val Gln Asp Gln Leu Gln Val Glu Arg Leu Gln His Thr
106 370 375 380

```

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```

107 Tyr Val Glu Ala Leu His Ala Tyr Val Ser Ile His His Pro His Asp
108 385 390 395 400
109 Arg Leu Met Phe Pro Arg Met Leu Met Lys Leu Val Ser Leu Arg Thr
110 405 410 415
111 Leu Ser Ser Val His Ser Glu Gln Val Phe Ala Leu Arg Leu Gln Asp
112 420 425 430
113 Lys Lys Leu Pro Pro Leu Leu Ser Glu Ile Trp Asp Val His Glu
114 435 440 445
117 <210> SEQ ID NO: 3
118 <211> LENGTH: 1383
119 <212> TYPE: DNA
120 <213> ORGANISM: Homo sapien
122 <400> SEQUENCE: 3
123 atgtcctctc ctaccacgag ttccctggat acccccctgc ctggaaatgg cccccctcag 60
124 cctggcgccc cttcttcttc acccaactgta aaggaggagg gtccggagcc gtggcccg 120
125 ggtccggacc ctgatgtccc aggcactgat gaggccagct cagcctgcag cacagactgg 180
126 gtcaccccag atcccgaaga ggaaccagag cgcaagcgaa agaagggcc agccccgaag 240
127 atgctgggcc acgagctttg ccgtgtctgt ggggacaagg cctccggctt ccactacaac 300
128 gtgctcagct gcgaaggctg caagggtctt tccggcgca gtgtggtccg tgggtggggc 360
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130 tgccagcagt gccggtgctg caagtgcagg gaggcaggga tgaggagca gtgcgtcctt 480
131 tctgaagaac agatccggaa gaagaagatt cggaaacagc agcaggagtc acagtcacag 540
132 tcgcagtcac ctgtggggcc gcagggcagc agcagctcag cctctggggc tggggcttcc 600
133 cctggtggat ctgaggcagg cagccagggc tccggggag gcgagggtgt ccagctaaca 660
134 gcggtccaag aactaatgat ccagcagttg gtggcggcc aactgcagt caacaaacgc 720
135 tccttctcgg accagcccaa agtcacgccc tggcccctgg gcgcagacc ccagtcgccg 780
136 gatccccgcc agcaacgctt tgcccacttc acggagctgg ccatcatctc agtcaggag 840
137 atcgtggact tcgctaagca agtgctggt ttctgcagc tgggcccggg ggaccagatc 900
138 gccctcctga aggcattcac tatcgagatc atgctgctag agacagccag gcgctacaac 960
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141 cggctgggcc tggacgacgc tgagtacgcc ctgctcatcg ccatcaacat cttctcggcc 1140
142 gaccggccca acgtgcagga gccgggccgc gtggaggcgt tgcagcagcc ctacgtggag 1200
143 gcgctgctgt cctacacgcg catcaagagg ccgcaggacc agctgcgctt cccgcgcag 1260
144 ctcatgaagc tggtagcct gcgcacgct agctctgtgc actcggagca ggtcttcgcc 1320
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146 tga 1383
148 <210> SEQ ID NO: 4
149 <211> LENGTH: 460
150 <212> TYPE: PRT
151 <213> ORGANISM: Homo sapien
153 <400> SEQUENCE: 4
154 Met Ser Ser Pro Thr Thr Ser Ser Leu Asp Thr Pro Leu Pro Gly Asn
155 1 5 10 15
156 Gly Pro Pro Gln Pro Gly Ala Pro Ser Ser Pro Thr Val Lys Glu
157 20 25 30
158 Glu Gly Pro Glu Pro Trp Pro Gly Gly Pro Asp Pro Asp Val Pro Gly
159 35 40 45
160 Thr Asp Glu Ala Ser Ser Ala Cys Ser Thr Asp Trp Val Ile Pro Asp

```



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210	Pro	Pro	Leu	Leu	Ser	Glu	Ile	Trp	Asp	Val	His	Glu
211		450					455					460

VERIFICATION SUMMARY

DATE: 01/30/2006

PATENT APPLICATION: US/10/565,495

TIME: 15:09:23

Input Set : A:\PU60406SEQLIST.txt

Output Set: N:\CRF4\01302006\J565495.raw

L:13 M:270 C: Current Application Number differs, Replaced Current Application Number

L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date